Docket No.: 13173-00025-US

AMENDMENTS TO THE CLAIMS

Listing of claims:

- 1. (Currently amended) A process for the preparation of a ketocarotenoid by culturing a genetically modified, nonhuman organism, which in comparison with the wild-type has a modified ketolase activity and a modified β -cyclase activity, and the modified β -cyclase activity is caused by a β -cyclase comprising the amino acid sequence SEQ. ID. NO. 2 SEQ ID NO: 2 or a sequence derived from this sequence by substitution, insertion or deletion of amino acids, which has an identity of at least 70% at the amino acid level with the sequence SEQ. ID. NO. 2 SEQ ID NO: 2.
- 2. (Original) The process according to claim 1, wherein a nonhuman organism is used which, as the wild-type, already has a ketolase activity, and the genetic modification causes an increase in the ketolase activity in comparison with the wild-type.
- 3. (Currently amended) The process according to claim 2, wherein for increasing the ketolase activity the gene expression of a nucleic acid encoding a ketolase is increased compared to the wild-type.
- 4. (Original) The process according to claim 3, wherein for increasing the gene expression a nucleic acid which encodes a ketolase is inserted into the organism.
- 5. (Original) The process according to claim 4, wherein, as a nucleic acid encoding a ketolase, a nucleic acid is inserted which encodes a ketolase comprising the amino acid sequence SEQ ID NO: 4 or a sequence derived from this sequence by substitution, insertion or deletion of amino acids, which has an identity of at least 70% at the amino acid level with the sequence SEQ ID NO: 4.
- 6. (Original) The process according to claim 1, wherein a nonhuman organism is used which, as the wild-type, has no ketolase activity and the genetic modification causes a ketolase activity in comparison with the wild-type.

- 7. (Original) The process according to claim 6, wherein a genetically modified organism is used which transgenically expresses a ketolase.
- 8. (Currently amended) The process according to claim 6 or 7, wherein, for causing the gene expression, a nucleic acid which encodes a ketolase is inserted into the organism.
- 9. (Currently amended) The process according to claim 8, wherein a nucleic acid is inserted encoding a ketolase comprising the amino acid sequence SEQ. ID. NO. 4 of SEQ ID NO: 4 or a sequence derived from this sequence by substitution, insertion or deletion of amino acids, which has an identity of at least 70% at the amino acid level with the sequence SEQ. ID. NO. 4 of SEQ ID NO: 4.

Claims 10-62 (Canceled).

- 63. (Currently amended) A genetically modified, nonhuman organism, where the genetic modification,
- A for the case where the wild-type organism already has a ketolase activity, increases the activity of a ketolase compared to the wild-type and
- B for the case where the wild-type organism has no ketolase activity, causes the activity of a ketolase compared to the wild-type,

and where the genetic modification,

- C for the case where the wild-type organism already has a β -cyclase activity, increases the activity of a β -cyclase compared to the wild-type and
- D for the case where the wild-type organism has no β -cyclase activity, causes the activity of a β -cyclase compared to the wild-type

and the β -cyclase activity increased according to C or caused according to D is caused by a β -cyclase comprising the amino acid sequence SEQ. ID. NO. 2 of SEQ ID NO: 2 or a sequence derived from this sequence by substitution, insertion or deletion of amino acids, which has an

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identity of at least 70% at the amino acid level with the sequence SEQ. ID. NO. 2 of SEQ ID NO: 2.

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Claims 64-71 (Canceled).

- 72. (New) The process according to claim 7, wherein, for causing the gene expression, a nucleic acid which encodes a ketolase is inserted into the organism.
- 73. (New) The process according to claim 72, wherein a nucleic acid is inserted encoding a ketolase comprising the amino acid sequence of SEQ ID NO: 4 or a sequence derived from this sequence by substitution, insertion or deletion of amino acids, which has an identity of at least 70% at the amino acid level with the sequence of SEQ ID NO: 4.